

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/053,526
Source:	OIPE
Date Processed by STIC:	5/7/02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- 3. Hand Carry directly to:
 - U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



Does Not Comply Corrected Diskette Needed

OIPE

Errors on P. 3

RAW SEQUENCE LISTING

3 <110> APPLICANT: INSTITUT CURIE

PATENT APPLICATION: US/10/053,526

DATE: 05/07/2002 TIME: 13:00:38

Input Set : A:\Sequence Listing

Output Set: N:\CRF3\05072002\J053526.raw

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CENTRE NATIONAL DE LA RECHERCHE SCIENTIFIQUE
      4
      5
              MUSEUM NATIONAL D'HISTOIRE NATURELLE
              INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE M
      6
      8 <120> TITLE OF INVENTION: Methods and compositions for effecting homologous
      9
              recombination
     11 <130> FILE REFERENCE: BET 01/1333
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/053,526
C--> 14 <141> CURRENT FILING DATE: 2002-04-18
     16 <150> PRIOR APPLICATION NUMBER: PCT/IB01/00749
     17 <151> PRIOR FILING DATE: 2001-05-03
     19 <150> PRIOR APPLICATION NUMBER: EP 00401218.3
     20 <151> PRIOR FILING DATE: 2000-05-03
     22 <160> NUMBER OF SEQ ID NOS: 11
     24 <170> SOFTWARE: PatentIn Ver. 2.1
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     28 <212> TYPE: DNA
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     32 <223> OTHER INFORMATION: Description of Artificial Sequence:
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                                                                           12
     65 <210> SEQ ID NO: 4
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RAW SEQUENCE LISTING DATE: 05/07/2002 PATENT APPLICATION: US/10/053,526 TIME: 13:00:38

Input Set : A:\Sequence Listing

Output Set: N:\CRF3\05072002\J053526.raw

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135 <220> FEATURE:

RAW SEQUENCE LISTING DATE: 05/07/2002 PATENT APPLICATION: US/10/053,526 TIME: 13:00:38

Input Set : A:\Sequence Listing

Output Set: N:\CRF3\05072002\J053526.raw

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     150
               oligonucleotide
                                                      requires caplonation for n - See
     152 <400> SEQUENCE: 10
W--> 153 nnnnntntnn ngnggng
     156 <210> SEQ ID NO: 11
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     158 <212> TYPE: DNA
     159 <213> ORGANISM: Artificial Sequence
     161 <220> FEATURE:
     162 <223> OTHER INFORMATION: Description of Artificial Sequence:
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/053,526

DATE: 05/07/2002 TIME: 13:00:39

Input Set : A:\Sequence Listing

Output Set: N:\CRF3\05072002\J053526.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:10; N Pos. 1,2,3,4,5,7,9,10,11,13,16

Seq#:11; N Pos. 1,2,3,5,7,9,11

VARIABLE LOCATION SUMMARY DATE: 05/07/2002 PATENT APPLICATION: US/10/053,526 TIME: 13:00:39

Input Set : A:\Sequence Listing

Output Set: N:\CRF3\05072002\J053526.raw

Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of $\langle 220 \rangle$ to $\langle 223 \rangle$ is MANDATORY if n's or Xaa's are present. in $\langle 220 \rangle$ to $\langle 223 \rangle$ section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:10; N Pos. 1,2,3,4,5,7,9,10,11,13,16

Seq#:11; N Pos. 1,2,3,5,7,9,11

VERIFICATION SUMMARY

DATE: 05/07/2002 TIME: 13:00:39 PATENT APPLICATION: US/10/053,526

Input Set : A:\Sequence Listing

Output Set: N:\CRF3\05072002\J053526.raw

L:13 M:270 C: Current Application Number differs, Replaced Application Number

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:153 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:10 L:153 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:10 L:153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0

L:166 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:11 L:166 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:11

L:166 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0